WHAT IS CLAIMED IS:

- 1. (currently amended) A method of determining the level of susceptibility of a subject to an environmental toxin capable of detoxification by a glutathione Stransferases <u>pi</u> comprising:
 - a. determining a first amount of one or more said glutathione S-transferases pi present in a biological sample from the subject;
 - b. contacting the biological sample with the environmental toxin;
 - c. determining a second amount of one or more <u>said</u> glutathione S-transferases pi present in the sample;

wherein the second amount of ene or more said glutathione S-transferases pi being lower than or similar to the first amount of ene or more said glutathione S-transferases pi indicates the subject having a higher level of susceptibility than a subject having the second amount of ene or more said glutathione S-transferases pi higher than the first amount of ene or more said glutathione S-transferases pi being higher than the first amount of ene or more said glutathione S-transferases pi being higher than the first amount of ene or more said glutathione S-transferases pi indicates the subject having a lower level of susceptibility than a subject having a second amount of ene or more said glutathione S-transferases pi lower than or similar to the first amount of ene or more said glutathione S-transferases pi lower than or similar to the first amount of ene or more said glutathione S-transferases pi, wherein said environmental toxin is MPTP or a metabolite thereof.

- 2. (original) The method of claim 1 wherein the subject is a mammal.
- 3. (original) The method of claim 1 wherein the subject is a human.
- 4. (original) The method of claim 1 wherein the subject is a rodent.
- 5. (original) The method of claim 1 wherein the subject is a mouse.
- 6. (original) The method of claim 1 wherein the biological sample is selected from the group consisting of plasma, brain and urine.

- 7. (Canceled)
- 8. (Canceled)
- 9. (original) The method of claim 1 wherein the environmental toxin is a toxin wherein the toxin or a metabolite thereof interferes with Complex I respiration in a cell's electron transport chain.
- 10. (original) The method of claim 1 wherein the environmental toxin is a toxin structurally similar to MPTP.
- 11. (Canceled)
- 12. (Currently Amended) The method of claim 1 wherein determining said first and second amounts of one or more said glutathione S-transferases <u>pi</u> involve determining the amount of mRNA encoding said one or more said glutathione S-transferases <u>pi</u>.
- 13. (Withdrawn) The method of claim 1 wherein determining said first and second amounts of one or more glutathione S-transferases involve the step of determining said one or more glutathione S-transferases by the method selected from the group consisting of: radioimmunoassay, enzyme immunoassay and immunoflurometric immuno assay.
- 14. (Currently Amended) The method of claim 12 wherein determining the amount of mRNA encoding said one-or more said glutathione S-transferases pi is determined using a primer selected from the group consisting of: SEQ ID NOS. 5-27, and SEQ ID NOS: 29-31, 33-35, 37-39, 40-43, 45-47, 49-51, 53-55, 57-59, 61-63, 65-67 and 69-71.
- 15. (Withdrawn) The method of claim 1 wherein determining said first and second amounts of one or more of one or more glutathione S-transferases involves determining the level of enzymatic activity corresponding to one or more glutathione S-transferases.
- 16. (Currently Amended) A method of determining risk of developing Parkinson's disease in a human subject comprising:

- a. determining a first amount of one or more said glutathione S-transferases pi present in a biological sample from the subject;
- b. contacting the biological sample with the environmental toxin;
- c. determining a second amount of one or more <u>said</u> glutathione Stransferases pi present in the sample;

wherein the second amount of <u>one or more said</u> glutathione S-transferases <u>pi</u> being lower than or similar to the first amount of <u>one or more said</u> glutathione S-transferases <u>pi</u> indicates the subject having a higher level of risk of developing Parkinson's disease than a subject having the second amount of <u>one or more said</u> glutathione S-transferases <u>pi</u> higher than the first amount of <u>one or more said</u> glutathione S-transferases <u>pi</u>, and wherein the second amount of <u>one or more said</u> glutathione S-transferases <u>pi</u> being higher than the first amount of <u>one or more said</u> glutathione S-transferases <u>pi</u> indicates the subject having a lower level of risk of developing Parkinson's disease than a subject having a second amount of the <u>one or more said</u> glutathione S-transferases <u>pi</u> lower than or similar to the first amount of the <u>one or more said</u> glutathione S-transferases <u>pi</u> lower than or similar to the first amount of the <u>one or more said</u> glutathione S-transferases <u>pi</u> wherein said environmental toxin is MPTP or a metabolite thereof.

- 17. (original) The method of claim 16 wherein the biological sample is selected from the group consisting of plasma, brain and urine.
- 18. (Canceled)
- 19. (original) The method of claim 16 wherein the environmental toxin is a toxin or a metabolite thereof interferes with Complex I respiration in a cell's electron transport chain.
- 20. (original) The method of claim 16 wherein the environmental toxin is a toxin structurally similar to MPTP.
- 21. (Canceled)

- 22. (Currently Amended) The method of claim 16 wherein determining said first and second amounts of the one or more said glutathione S-transferases <u>pi</u> involve determining the amount of mRNA encoding said the one or more said glutathione S-transferases <u>pi</u>.
- 23. (Withdrawn) The method of claim 16 wherein determining said first and second amounts of the one or more glutathione S-transferases involve the step of determining said glutathione S-transferase by the method selected from the group consisting of: radioimmunoassay, enzyme immunoassay and immunoflurometric immunoassay.
- 24. (Withdrawn) The method of claim 23 wherein determining the amount of mRNA encoding said at least one glutathione S-transferases is determined using a primer selected from the group consisting of: SEQ ID NOS: 29-31, 33-35, 37-39, 41-43, 45-47, 49-51, 53-55, 57-59, 61-63, 65-67, and 69-71.
- 25. (Withdrawn) The method of claim 16 wherein determining said first and second amounts of the one or more glutathione S-transferases involves determining the level of enzymatic activity corresponding to the one or more glutathione S-transferases.
- 26. (withdrawn) A method to determine genetic susceptibility to an environmental toxin involving contacting a biological sample from a subject with a composition consisting essentially of one or more labeled probes each of which binds selectively to a region on mouse chromosome 1, from D1Mit113 to D1Mit293 under conditions in which one or more labeled probes form a stable hybridization complex with DNA in the region of mouse chromosome 1 from D1Mit113 to D1Mit293 and detecting the hybridization complex, wherein said hybridization complex is indicative of a genetic susceptibility to an environmental toxin.
- 27. (withdrawn) A method to determine the susceptibility of a subject to develop Parkinson's disease, the method comprising: contacting a biological sample from a subject with a composition consisting essentially of a plurality of labeled probes each of which selectively binds to a region of human Glutathione S-transferase pi within a segment of human GSTP1 (chr11:69874218-69877056); GST mu (hGSTm4:

chr1:110677671-110687021, hGSTm2: chr1:110689771-110696957, hGSTm3: chr1:110758175-110761973, hGSTm1:chr1:110709530-110715415; or GST alpha (hGSTa1:chr6:52658645-52670705, hGSTa2:chr6:52617245-52630341; hGSTa3:chr6:52763514-52776547, hGSTa4:chr6:52844814-52862163 under conditions which a plurality of labeled probes form stable hybridization complexes with a segment of human GSTP1 (chr11:69874218-69877056); GST mu (hGSTm4: chr1:110677671-110687021, hGSTm2: chr1:110689771-110696957, hGSTm3: chr1:110758175-110761973, hGSTm1:chr1:110709530-110715415; or GST alpha (hGSTa1:chr6:52658645-52670705, hGSTa2:chr6:52617245-52630341; hGSTa3:chr6:52763514-52776547, hGSTa4:chr6:52844814-52862163; and detecting the hybridization complex wherein said hybridization complex is indicative of increased susceptibility of a subject to develop Parkinson's disease.